

DMCI0099.ST25.txt
SEQUENCE LISTING

<110> Havkin-Frenkel, Daphna
Podstolski, Andrzej
Dixon, Richard A.

<120> Vanillin Biosynthetic Pathway Enzyme From Vanilla Planifolia

<130> DMCI0099

<150> 09/462,576
<151> 2000-05-22

<150> PCT/US98/14895
<151> 1998-07-15

<150> 60/052,604
<151> 1997-07-15

<150> 60/272,415
<151> 2001-02-28

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<170> PatentIn version 3.1

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145 150 155 160Ser Trp Thr Phe Ser Thr Thr Gly Ala Leu Glu Ala Ala Tyr Thr Gln
165 170 175Leu Thr Gly Ser Thr Leu Ser Glu Gln Gln Leu Val Asp Cys Ala Ser
180 185 190Ala Phe Asn Asn Phe Gly Cys Gly Gly Leu Pro Ser Gln Ala Phe Glu
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Leu Gly Val Met Gly Ile Cys Asn Phe Lys Gln Glu Asn Val Gly Val
225 230 235 240

Lys Val Ile Asp Ser Ile Asn Ile Thr Leu Gly Ala Glu Asp Glu Leu
245 250 255

Lys His Ala Val Gly Leu Val Arg Pro Val Ser Val Ala Phe Glu Val
260 265 270

Val Lys Gly Phe Asn Leu Tyr Lys Lys Gly Val Tyr Ser Ser Asp Thr
275 280 285

Cys Gly Arg Asp Pro Met Asp Val Asn His Ala Val Leu Ala Val Gly
290 295 300

Tyr Gly Val Glu Asp Gly Ile Pro Tyr Trp Leu Ile Lys Asn Ser Trp
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 35 40 45

Ile Leu Gly Gln Ser Arg His Val Leu Ser Phe Ala Arg Phe Thr His
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Arg Tyr Gly Lys Lys Tyr Gln Asn Val Glu Glu Met Lys Leu Arg Phe
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Ser Ile Phe Lys Glu Asn Leu Asp Leu Ile Arg Ser Thr Asn Lys Lys
85 90 95

Gly Leu Ser Tyr Lys Leu Gly Val Asn Gln Phe Ala Asp Leu Thr Trp
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Gln Glu Phe Gln Arg Thr Lys Leu Gly Ala Ala Gln Asn Cys Ser Ala
115 120 125

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Lys Asp Trp Arg Glu Asp Gly Ile Val Ser Pro Val Lys Asp Gln Gly
145 150 155 160

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165 170 175

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180 185 190

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195 200 205

Gly Gly Leu Pro Ser Gln Ala Phe Glu Tyr Ile Lys Ser Asn Gly Gly
210 215 220

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225 230 235 240

Lys Phe Ser Ala Glu Asn Val Gly Val Gln Val Leu Asn Ser Val Ser
245 250 255

Ile Thr Leu Gly Ala Glu Asp Glu Leu Lys His Ala Val Gly Leu Val
260 265 270

Arg Pro Val Ser Ile Ala Phe Glu Val Ile His Ser Phe Arg Leu Tyr
275 280 285

Lys Ser Gly Val Tyr Thr Asp Ser His Cys Gly Ser Thr Pro Met Asp
290 295 300

Val Asn His Ala Val Leu Ala Val Gly Tyr Gly Val Glu Asp Gly Val
305 310 315 320

Pro Tyr Trp Leu Ile Lys Asn Ser Trp Gly Ala Asp Trp Gly Asp Lys
325 330 335

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 35 40 45

Leu Gly Ala Leu Gly Arg Thr Arg His Ala Leu Arg Phe Ala Arg Phe
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Ala Val Arg Tyr Gly Lys Ser Tyr Glu Ser Ala Ala Glu Val Arg Arg
 65 70 75 80

Arg Phe Arg Ile Phe Ser Glu Ser Leu Glu Glu Val Arg Ser Thr Asn
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Arg Lys Gly Leu Pro Tyr Arg Leu Gly Ile Asn Arg Phe Ser Asp Met
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 115 120 125

Ser Ala Thr Lys Gly Asn His Leu Met Arg Asp Ala Ala Ala Leu Pro
 130 135 140

Glu Thr Lys Asp Trp Arg Glu Asp Gly Ile Val Ser Pro Val Lys Asn
 145 150 155 160

Gln Ala His Cys Gly Ser Cys Trp Thr Phe Ser Thr Thr Gly Ala Leu
 165 170 175

Glu Ala Ala Tyr Thr Gln Ala Thr Gly Lys Asn Ile Ser Leu Ser Glu
 180 185 190

Gln Gln Leu Val Asp Cys Ala Gly Gly Phe Asn Asn Phe Gly Cys Asn
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195

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Gly Gly Leu Pro Ser Gln Ala Phe Glu Tyr Ile Lys Tyr Asn Gly Gly
 210 215 220

Ile Asp Thr Glu Glu Ser Tyr Pro Tyr Lys Gly Val Asn Gly Val Cys
 225 230 235 240

His Tyr Lys Ala Glu Asn Ala Ala Val Gln Val Leu Asp Ser Val Asn
 245 250 255

Ile Thr Leu Asn Ala Glu Asp Glu Leu Lys Asn Ala Val Gly Leu Val
 260 265 270

Arg Pro Val Ser Val Ala Ala Phe Gln Val Ile Asp Gly Phe Arg Gln
 275 280 285

Tyr Lys Ser Gly Val Tyr Thr Ser Asp His Cys Gly Thr Thr Pro Asp
 290 295 300

Asp Val Asn His Ala Val Leu Ala Val Gly Tyr Gly Val Glu Asn Gly
 305 310 315 320

Val Pro Tyr Trp Leu Ile Lys Asn Ser Trp Gly Ala Asp Trp Gly Asp
 325 330 335

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 35 40 45

Leu Gln Val Val Gly Lys Thr Arg His Ala Leu Ser Phe Ala Arg Phe
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Ala His Arg Tyr Gly Lys Arg Tyr Glu Ser Val Glu Glu Ile Lys Gln
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Arg Phe Glu Val Phe Leu Asp Asn Leu Lys Met Ile Arg Ser His Asn
85 90 95

Lys Lys Gly Leu Ser Tyr Lys Leu Gly Val Asn Glu Phe Thr Asp Leu
100 105 110

Thr Trp Asp Glu Phe Arg Arg Asp Arg Leu Gly Ala Ala Gln Asn Cys
115 120 125

Ser Ala Thr Thr Lys Gly Asn Leu Lys Val Thr Asn Val Val Leu Pro
130 135 140

Glu Thr Lys Asp Trp Arg Glu Ala Gly Ile Val Ser Pro Val Lys Asn
145 150 155 160

Gln Gly Lys Cys Gly Ser Cys Trp Thr Phe Ser Thr Thr Gly Ala Leu
165 170 175

Glu Ala Ala Tyr Ser Gln Ala Phe Gly Lys Gly Ile Ser Leu Ser Glu
180 185 190

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195 200 205

Gly Gly Leu Pro Ser Gln Ala Phe Glu Tyr Ile Lys Ser Asn Gly Gly
210 215 220

Leu Asp Thr Glu Glu Ala Tyr Pro Tyr Thr Gly Lys Asn Gly Leu Cys
225 230 235 240

Lys Phe Ser Ser Glu Asn Val Gly Val Lys Val Ile Asp Ser Val Asn
245 250 255

Ile Thr Leu Gly Ala Glu Asp Glu Leu Lys Tyr Ala Val Ala Leu Val
260 265 270

Arg Pro Val Ser Ile Ala Phe Glu Val Ile Lys Gly Phe Lys Gln Tyr
275 280 285

Lys Ser Gly Val Tyr Thr Ser Thr Glu Cys Gly Asn Thr Pro Met Asp
290 295 300

Val Asn His Ala Val Leu Ala Val Gly Tyr Gly Val Glu Asp Gly Val
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Ala Leu Gly Arg Thr Arg Asp Ala Leu Arg Phe Ala Arg Phe Ala Val
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65 70 75 80

Arg Ile Phe Ser Glu Ser Leu Gln Leu Val Arg Ser Thr Asn Arg Lys
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Gly Leu Ser Tyr Arg Leu Gly Ile Asn Arg Phe Ala Asp Met Ser Trp
100 105 110

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115 120 125

Thr Leu Thr Gly Asn His Arg Met Arg Ala Ala Ala Val Ala Leu Pro
130 135 140

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145 150 155 160

Gln Gly His Cys Gly Ser Cys Trp Thr Phe Ser Thr Thr Gly Ala Leu
165 170 175

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Glu Ala Ala Tyr Thr Gln Ala Thr Gly Lys Pro Ile Ser Leu Ser Glu
180 185 190

Gln Gln Leu Val Asp Cys Gly Leu Ala Phe Asn Asn Phe Gly Cys Asn
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Gly Gly Leu Pro Ser Gln Ala Phe Glu Tyr Ile Lys Tyr Asn Gly Gly
210 215 220

Leu Asp Thr Glu Glu Ser Tyr Pro Tyr Gln Gly Val Asn Gly Ile Ser
225 230 235 240

Lys Phe Lys Asn Glu Asn Val Gly Val Lys Val Leu Asp Ser Val Asn
245 250 255

Ile Thr Leu Gly Ala Glu Asp Glu Leu Lys Asp Ala Val Gly Leu Val
260 265 270

Arg Pro Val Ser Val Ala Phe Glu Val Ile Thr Gly Phe Arg Leu Tyr
275 280 285

Lys Ser Gly Val Val Thr Ser Asp His Cys Gly Thr Thr Pro Met Asp
290 295 300

Val Asn His Ala Val Leu Ala Val Gly Tyr Gly Val Glu Asp Gly Val
305 310 315 320

Pro Tyr Trp Leu Ile Lys Asn Ser Trp Gly Ala Asp Trp Gly Asp Glu
325 330 335

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Cys Ala Ser Tyr Pro Ile Val Ala
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Pro Ile Arg Pro Val Thr Asp Arg Ala Ala Ser Ala Leu Glu Ser Thr
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Val Phe Ala Ala Leu Gly Arg Thr Arg Asp Ala Leu Arg Phe Ala Arg
50 55 60

Phe Ala Val Arg Tyr Gly Lys Ser Tyr Glu Ser Ala Ala Glu Val His
65 70 75 80

Lys Arg Phe Arg Ile Phe Ser Glu Ser Leu Gln Leu Val Arg Ser Thr
85 90 95

Asn Arg Lys Gly Leu Ser Tyr Arg Leu Gly Tyr Asn Arg Phe Ala Asp
100 105 110

Met Ser Trp Glu Glu Phe Arg Ala Thr Arg Leu Gly Ala Ala Gln Asn
115 120 125

Cys Ser Ala Thr Leu Thr Gly Asn His Arg Met Arg Ala Ala Ala Val
130 135 140

Ala Leu Pro Glu Thr Lys Asp Trp Arg Glu Asp Gly Ile Val Ser Pro
145 150 155 160

Val Lys Asn Gln Gly His Cys Gly Ser Cys Trp Thr Phe Ser Thr Thr
165 170 175

Gly Ala Leu Glu Ala Ala Tyr Thr Gln Ala Thr Gly Lys Pro Ile Ser
180 185 190

Leu Ser Glu Gln Gln Leu Val Asp Cys Gly Phe Ala Phe Asn Asn Phe
195 200 205

Gly Cys Asn Gly Gly Leu Pro Ser Gln Ala Phe Glu Tyr Ile Lys Tyr
210 215 220

Asn Gly Gly Leu Asp Thr Glu Glu Ser Tyr Pro Tyr Gln Gly Val Asn
225 230 235 240

Gly Ile Cys Lys Phe Lys Asn Glu Asn Val Gly Val Lys Val Leu Asp
245 250 255

Ser Val Asn Ile Thr Leu Gly Ala Glu Asp Glu Leu Lys Asp Ala Val
260 265 270

Gly Leu Val Arg Pro Val Ser Val Ala Phe Glu Val Ile Thr Gly Phe
275 280 285

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Arg Leu Tyr Lys Ser Gly Val Tyr Thr Ser Asp His Cys Gly Thr Thr
290 295 300

Pro Met Asp Val Asn His Ala Val Leu Ala Val Gly Tyr Gly Val Glu
305 310 315 320

Asp Gly Val Pro Tyr Trp Leu Ile Lys Asn Ser Trp Gly Ala Asp Trp
325 330 335

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Pro Ile Arg Ser Val Thr Asp His Ala Ala Ser Ala Leu Glu Ser Thr
35 40 45

Val Ile Ala Ala Leu Gly Arg Thr Arg Gly Ala Leu Arg Phe Ala Arg
50 55 60

Phe Ala Val Arg Gly His Lys Arg Tyr Gly Asp Ala Ala Glu Val Gln
65 70 75 80

Arg Arg Phe Arg Ile Phe Ser Glu Ser Leu Glu Leu Val Arg Ser Thr
85 90 95

Asn Arg Arg Gly Leu Pro Tyr Arg Leu Gly Ile Asn Arg Phe Ala Asp
100 105 110

Met Ser Trp Glu Glu Phe Gln Ala Ser Arg Leu Gly Ala Ala Gln Asn
115 120 125

Cys Ser Ala Thr Leu Ala Gly Asn His Arg Met Arg Asp Ala Pro Ala
130 135 140

Leu Pro Glu Thr Lys Asp Trp Arg Glu Asp Gly Ile Val Ser Pro Val
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145 150 160

Lys Asp Gln Gly His Cys Gly Ser Cys Trp Pro Phe Ser Thr Thr Gly
165 170 175

Ser Leu Glu Ala Arg Tyr Thr Gln Ala Thr Gly Pro Pro Val Ser Leu
180 185 190

Ser Glu Gln Gln Leu Ala Asp Cys Ala Thr Arg Tyr Asn Asn Phe Gly
195 200 205

Cys Ser Gly Gly Leu Pro Ser Gln Ala Phe Glu Tyr Ile Lys Tyr Asn
210 215 220

Gly Gly Leu Asp Thr Glu Glu Ala Tyr Pro Tyr Thr Gly Val Asn Gly
225 230 235 240

Ile Cys His Tyr Lys Pro Glu Asn Ala Gly His Lys Val Leu Asp Ser
245 250 255

Val Asn Ile Thr Leu Val Ala Glu Asp Glu Leu Lys Asn Ala Val Gly
260 265 270

Leu Val Arg Pro Val Ser Val Ala Phe Gln Val Ile Asn Gly Phe Arg
275 280 285

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290 295 300

Met Asp Val Asn His Ala Val Leu Ala Val Gly Tyr Gly Val Glu Asn
305 310 315 320

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325 330 335

Asp Asn Gly Tyr Phe Thr Met Glu Met Gly Lys Asn Met Cys Gly Ile
340 345 350

Ala Thr Cys Ala Ser Tyr Pro Ile Val Ala
355 360

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